

Figure 1

Keratinocyte Derived Interferon

1 CCA CGC GTC CGG GAT TTT TTA GCT TGC AAA AAA AAT GAG CAC CAA ACC TGA TAT GAT TCA 60
1 M S T K P D M I Q 9

61 AAA GTG TTT GTG GCT TGA GAT CCT TAT GGG TAT ATT CAT TGC TGG CAC CCT ATC CCT GGA 120
9 K C L W L E I L M G I F I A G T L S L D 29

121 CTG TAA CTT ACT GAA CGT TCA CCT GAG AAG AGT CAC CTG GCA AAA TCT GAG ACA TCT GAG 180
29 C N L L N V H L R R V T W Q N L R H L S 49

181 TAG TAT GAG CAA TTC ATT TCC TGT AGA ATG TCT ACG AGA AAA CAT AGC TTT TGA GTT GCC 240
49 S M S N S F P V E C L R E N I A F E L P 69

241 CCA AGA GTT TCT GCA ATA CAC CCA ACC TAT GAA GAG GGA CAT CAA GAA GGC CTT CTA TGA 300
69 Q E F L Q Y T Q P M K R D I K K A F Y E 89

301 AAT GTC CCT ACA GGC CTT CAA CAT CTT CAG CCA ACA CAC CTT CAA ATA TTG GAA AGA GAG 360
89 M S L Q A F N I F S Q H T F K Y W K E R 109

361 ACA CCT CAA ACA AAT CCA AAT AGG ACT TGA TCA GCA AGC AGA GTA CCT GAA CCA ATG CTT 420
109 H L K Q I Q I G L D Q Q A E Y L N Q C L 129

421 GGA GGA AGA CGA GAA TGA AAA TGA AGA CAT GAA AGA AAT GAA AGA GAA TGA GAT GAA ACC 480
129 E E D E N E N E D M K E M K E N E M K P 149

481 CTC AGA AGC CAG GGT CCC CCA GCT GAG CAG CCT GGA ACT GAG GAG ATA TTT CCA CAG GAT 540
149 S E A R V P Q L S S L E L R R Y F H R I 169

541 AGA CAA TTT CCT GAA AGA AAA GAA ATA CAG TGA CTG TGC CTG GGA GAT TGT CCG AGT GGA 600
169 D N F L K E K K Y S D C A W E I V R V E 189

601 AAT CAG AAG ATG TTT GTA TTA CTT TTA CAA ATT TAC AGC TCT ATT CAG GAG GAA ATA AGA 660
189 I R R C L Y Y F Y K F T A L F R R K * 207

661 ATC ATC TAC CTT CAA GCA AGA ATT AAC AGA GAT TGT GGC TAC GCA AAT GCA CCT AAA AAG 720
721 GGT GAA ATA TAT CTG AAA TGT ACC TGG TTC TGC CCT TGG AAG CCA CTT OCT CTT CAT GCC 780
781 ACT AAC AGC ATG CTG CCA AAC TGT TCA GAT TCA AGA TTA TTC CAA GCG CAG GGC CCA AAT 840
841 GTT ATA GCC AAA GAA AGT CTT ATG ATA AAA GTG AGG CAA ATT TCA GCC AAG AAG TTA GAA 900
901 GAG ATG TTT AAA AGA ACA AGA ACA AAT TGT GGA TCA TGG TAT ATG CAG GCT ATC AGC AGA 960
961 AGG ATC AGA CAA TAA AAT GAG TTA GTG CAA ACC ATT TAG TAA AAA TAA CTA TCA GCA GAG 1020
1021 TTG TTC CAG ATT AAA AAT AGT ACT ACA AGC TTG TAA AGG AGT TAG GAC ATG CAA GCT ACT 1080
1081 GAG CAT AAA ATA TAT ACT TGC TAT TTT TCA TGA CTT TCT CTA ATA AAG TCT TTG ACT GTT 1140
1141 CTC TCT AAT AAA AAA AAA AAA AAA AAA AAA 1170

CCF 11220 20555000

Figure 2

HKAPI15	10	KCLWLEILMGIFIAG...TSLDCNLLNVH..LRRVTWQNLRHLSSMSNS	54
		.. ::: ::::: . :: . . : :::	
INF-omega	1	MALLFPLLAALVMTSYSVPGSLGCDLPQNHGLLSRNTLVLLHQMRRIS..	48
HKAPI15	55	FPVECLRENIAFELPQEFQYTQPMKRDIKKAFYEMSLQAFNIF.SQHTF	103
		. ::: . : ::: : . . .	
INF-omega	49	.PFLCLKDRRDFRFPQEMVKGSQLOKAHVMSVLHEMLQQIFSLFHTERS	97
HKAPI15	104	KYWKERHLKQIQIGLDQQAAYLNQCLEEDENENEDMKEMKENEMKPSEAR	153
		. . ::: . ::: :	
INF-omega	98	AAWNMTLLDQLHTELHQQLQHLETCLLQVVGESESAGAISS.....	138
HKAPI15	154	VPQLSSLELRRYFHRIDNFLKEKKYSDCAWEIVRVEIRRCLYYFYKFTAL	203
		.. . : : : : :	
INF-omega	139PALTLRRYFQGIRVYLKEKKYSDCAWEVVRMEIMKSLFLSTNMQER	184
HKAPI15	204	FRRK	207
		: .	
INF-omega	185	LRSK	188

Figure 3

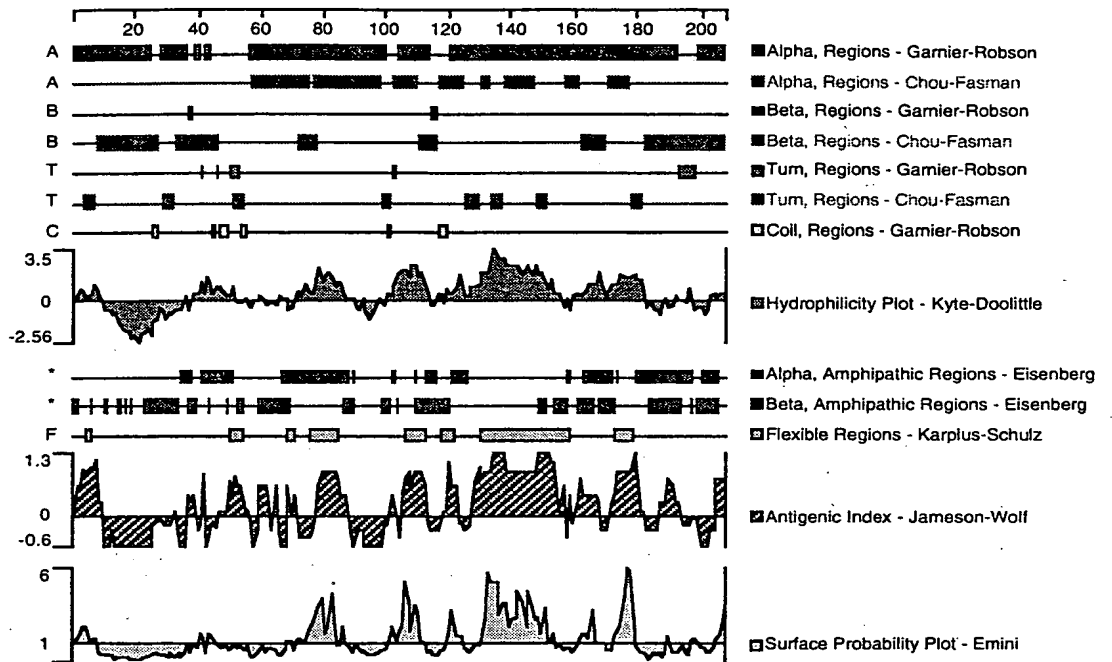


FIGURE 4
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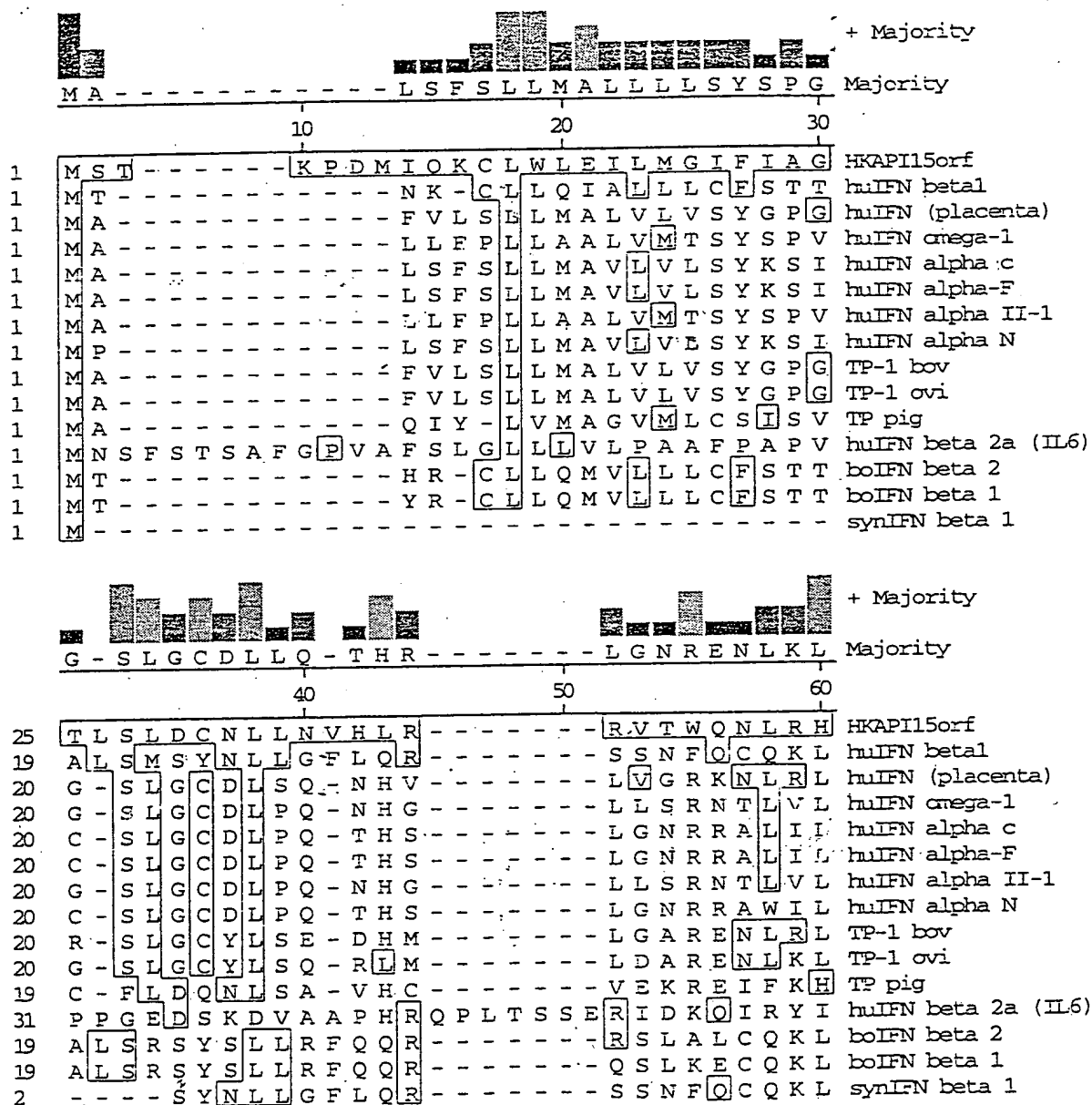


FIGURE 4
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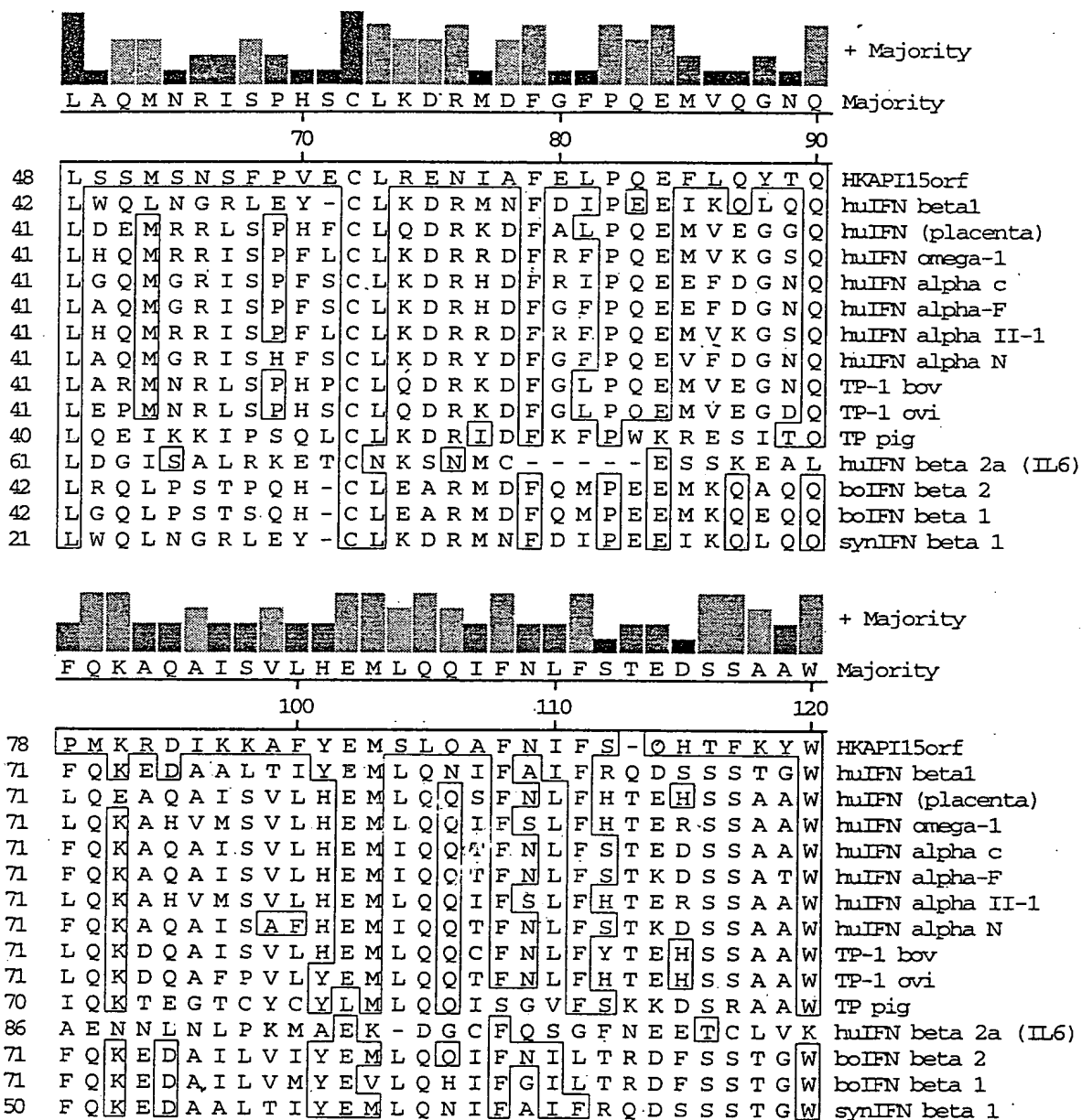


FIGURE 4
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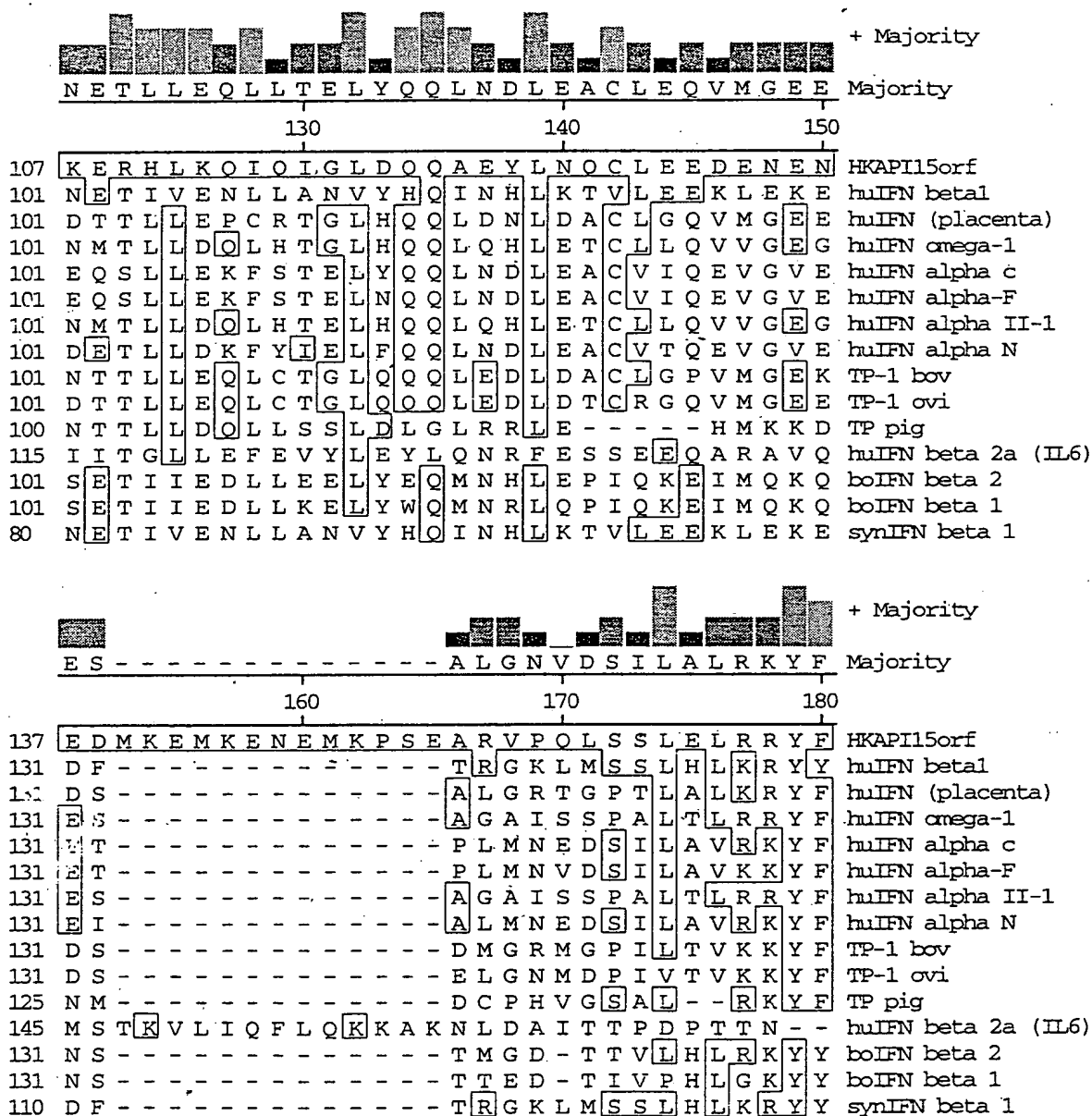


FIGURE 4
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